TRANSDUCTIONAL ANALYSIS OF MOMOPHASIC TYPES OF SALMONELLA

(Report by Tetsuo line and Joshua Lederberg*)

The specificity of flagellar (H) antigen in Salmonella is controlled by two distinction loci, phase-1 by \underline{H}_1 and phase-2 by \underline{H}_2 . Which one is manifested in a given clone depends on the phase determinant at the \underline{H}_2 locus. That is, the alternation of \underline{H}_2 state leads to the alternative expression, which has been known as phase variation of H-antigen (the Annual Report, 1956).

Some Selmonella strains do not express two phases but only one. Those strains are called monophasic-1 or -2 strains depending on their fixed phase, either phase-1 or phase-2 respectively. Three additional groups of the genes which are involved in the production of H antigen were disclosed by transductional analysis of the monophasic strains.

S. abortus—equi CDC-26 is stable in phase-2, enx-type.

A rere alternative phase can occasionally be obtained by antiserum selection, resulting in a equally stable phase-1 (a).

Transductions were carried out from enx-phase of CDC-26 to i-phase
of a diphasic strain of S. typhimurium, TM2 i:1.2 (such transduction
is designated CDC-26 (a):enx -x TM2 i:1.2). Among 65 transductional
clones which had been selected on semisolid nutrient gelatine-agar
media (NGA), 4 expressed diphasic a:1.2 type, 42 diphasic i:enx and
19 monophasic-2 enx which carry a hidden H₁. Thus, when a is
transduced a fraction of the transductional clones become monophasic.

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the resulting transductions remain diphasic, whereas when eax is transduced a fraction of the transductional clones become monophasic. By anti-eax selection, i-phase cultures were obtained from the monophasic eax-transductional clones. The i-phase cultures obtained are also monophasic. The stabilization of the H₂ state in S. abortine-equi is therefore caused by a factor which is linked to H₂. The factor will be given a symbol Vh₂.

S. typhimurium Sw1061 is a monophasic-2 mutant of a diphasic strain TM2 i:1.2. The culture reacts to anti-1.2 serum but not to anti-1 serum. However, the strain frequently produces nonmotile H-negative (non-flagellar) subclones, which in turn revert to motile cells with 1.2 antigen in successive cultures.

From the transduction, diphasic S. abony CDC-103 b:enx ---x Sw1061, monophasic-2 enx, diphasic b:1.2 and a small number of i:1.2 types were obtained. The change from the monophasic type to diphasic types was always coupled with the loss of the ability to oscillate between motile and non-motile types.

Thete results are consistent with the following explanation.

In Sw1061, H₁ is inactive; on the other hand H₂ changes its state as in usual diphasic strains. Consequently, when H₂ is active, phase-2 antigen, 1.2, is produced, and when H₂ changes to inactive, that is both H₁ and H₂ are inactive, the cell

cannot manifest produce H antigen and become non-motile. The production of diphasic | i:1.2 type suggests that the inactivation of H1 is not caused by an intrinsic change of $\underline{\underline{H}}_1$ itself but by an inhibition of its function by a gene linked to H1. The linkage and the recombination between $\underline{\underline{H}}_1$ and $\underline{\underline{H}}_1$ -inhibituractivity controller were confirmed on $\underline{\text{Fla}}_1^+$ (linked to $\underline{\text{H}}_1$) transductions from SW1061 --- x S. heidelberg Flag r:1.2, from which monophasio-2 -(i):1.2 and diphasic i:1.2 as well as diphasic r:1.2 were obtained. The H_-activity controller is designated Ah_ was discovered in two other monophasic-2 strains of S. typhimurium, SW629 and SW547. All of three Ah are non-vallelic and linked to H1 and Flat. The parrel type of gene, Ah , which inhibites the function of $\underline{\mathbb{H}}_2$ was disclosed on four monophasic-1 strains of \underline{S} , typhimurium . MARKANIAN AND ALL of them are linked to H2. Both Ab1 and Ah are phase specific but are not concerned with the specificity of antigen types, which are determined exclusively by $\underline{\mathbb{H}}_1$ and $\underline{\mathbb{H}}_2$. (the detail will be reported in) }